



#7

<110> Coleman, Timothy
Mansfield, Brian

<120> Scaffold Fusion Polypeptides, Composition for Making the Same,
and Methods of Using the Same.

<130> PF537

<140> us/10/057,890

<141> 2002-01-29

<150> 60/265,782

<151> 2001-01-31

<150> 60/265,858

<151> 2001-01-31

<160> 32

<210> 1

<211> 8

<212> PRT

<213> Homo sapiens

<400> 1

Glu Phe Leu Ile Val Ile Lys Ser

1

5

<210> 2

<211> 27

<212> PRT

<213> Homo sapiens

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<222> (1)..(1)

<223> Xaa equals either the naturally occurring L-amino acid Phe or Tyr

<220>

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<222> (2)..(2)

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<222> (4)..(7)

<223> Xaa equals any of the naturally occurring L-amino acids, one or
two residues may be missing

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<222> (9)..(9)

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<222> (10)..(10)
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 <222> (22)..(26)
 <223> Xaa equals any of the naturally occurring L-amino acids, one or
 two residues may be missing.

<400> 2
 Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Phe Xaa Xaa Xaa Xaa Xaa Leu
 1 5 10 15
 Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His
 20 25

<210> 3

<211> 10
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (5)..(5)
<223> Xaa equals any of the naturally occurring L-amino acids, one or two residues may be missing

<400> 3
Cys Pro Glu Cys Xaa His Gln Arg Thr His
1 5 10

<210> 4
<211> 8
<212> PRT
<213> Homo sapiens

<220>
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<222> (1)..(1)
<223> Xaa equals either the naturally occurring L-amino acid Phe or Tyr

<220>
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<220>
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<223> Xaa equals any of the naturally occurring L-amino acids, one or two residues may be missing

<400> 4
Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys
1 5

<210> 5
<211> 7
<212> PRT
<213> Homo sapiens

<220>
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<222> (2)..(6)
<223> Xaa equals any of the naturally occurring L-amino acids, one or two residues may be missing

<400> 5
His Xaa Xaa Xaa Xaa Xaa His
1 5

<210> 6
<211> 6
<212> PRT

a
Cont

<213> Homo sapiens

<400> 6

Tyr Lys Cys Gly Leu Cys
1 5

<210> 7

<211> 5

<212> PRT

<213> Homo sapiens

<400> 7

His Gln Arg Val His
1 5

<210> 8

<211> 7

<212> PRT

<213> Homo sapiens

<400> 8

Thr Gly Glu Lys Pro Tyr Lys
1 5

<210> 9

<211> 209

<212> PRT

<213> Homo sapiens

<400> 9

Met Asp Tyr Val Ser Ser Tyr Asp Asn Tyr Tyr Thr Ser Cys Lys Asn
1 5 10 15

Val Lys Ala Ala Arg Tyr Ser Val Gly Val Gly Asn Met Val Asn Cys
20 25 30

Lys Arg Lys Ser Met Thr Asp Tyr Asn Ala Ser Asp Thr Val Trp Ala
35 40 45

His Tyr Ala Ala Ala Trp Asp Gly Asn Thr Met Cys Thr Gly Tyr Gly
50 55 60

Ser Gly Thr Asp Arg Tyr Ala Val Val His Ala Val Ala Lys Ala Arg
65 70 75 80

Thr Val Thr Gly Val Val Thr Ser Val Thr Trp Val Val Ala Val Ala
85 90 95

Ser Gly Thr Arg Ser Lys Gly His Tyr Thr Cys Ser Ser His Tyr Ser
100 105 110

Tyr Trp Lys Asn Thr Lys Val Gly Val Val Met Val Cys Tyr Ser Gly
115 120 125

Lys Thr Arg Cys Arg Asn Lys Lys Arg His Arg Ala Val Arg Thr Met
130 135 140

a!
Cms

Val Tyr Trp Ala Tyr Asn Val Asn Thr Gly Asn Asn Cys Ser Ser Ser
145 150 155 160

Asn Arg Asp Ala Met Val Thr Thr Gly Met Thr His Cys Cys Asn Tyr
165 170 175

Ala Val Gly Lys Arg Asn Tyr Val Lys His Ala Lys Arg Cys Lys Cys
180 185 190

Cys Ser Ala Arg Ala Ser Ser Val Tyr Thr Arg Ser Thr Gly Ser Val
195 200 205

Gly

<210> 10

<211> 138

<212> PRT

<213> Homo sapiens

<400> 10

Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
1 5 10 15

Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Tyr Lys
20 25 30

Cys Gly Leu Cys Ala Ala Ala Gln Trp Asp Phe Gly Asn Thr Met Cys
35 40 45

Gln His Gln Arg Val His Gly His His His His Ser Tyr Lys Cys Gly
50 55 60

Leu Cys Thr Arg Ser Gln Lys Glu Gly Leu His Tyr Thr Cys Ser Ser
65 70 75 80

His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu
85 90 95

Lys Ile His Gln Arg Val His Gly Gly Gly Gly Ser Tyr Lys Cys Gly
100 105 110

Leu Cys Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser Asn Arg
115 120 125

Leu Asp Gly His Gln Arg Val His Ala Ala
130 135

<210> 11

<211> 30

<212> PRT

<213> Homo sapiens

<400> 11

Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
1 5 10 15

Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala
 20 25 30

<210> 12
 <211> 13
 <212> PRT
 <213> Homo sapiens

<400> 12
 Ala Ala Ala Gln Trp Asp Phe Gly Asn Thr Met Cys Gln
 1 5 10

<210> 13
 <211> 32
 <212> PRT
 <213> Homo sapiens

<400> 13
 Thr Arg Ser Gln Lys Glu Gly Leu His Tyr Thr Cys Ser Ser His Phe
 1 5 10 15

Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile
 20 25 30

<210> 14
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 14
 Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser Asn Arg Leu Asp
 1 5 10 15

Gly

<210> 15
 <211> 6
 <212> PRT
 <213> Homo sapiens

<400> 15
 Gly His His His His Ser
 1 5

<210> 16
 <211> 5
 <212> PRT
 <213> Homo sapiens

<400> 16
 Gly Gly Gly Gly Ser
 1 5

<210> 17
 <211> 72
 <212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotides used to join DNA fragments

<400> 17

gatctgtaat acgactcact atagggcacc atatggacta ccaggtttct tctccgatct 60
acgacatcaa ct 72

<210> 18

<211> 72

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotides used to join DNA fragments

<400> 18

gcagcgatct gtttaacggt gattttctgg cacgggtcag aggtgtagta gttgatgtcg 60
tagatcggag aa 72

<210> 19

<211> 72

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotides used to join DNA fragments

<400> 19

aatcaacggt aaacagatcg ctgcttaca atgcgggtctg tgcgctgctg ctcagtggga 60
cttcggtaac ac 72

<210> 20

<211> 79

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotides used to join DNA fragments

<400> 20

ggatccggat ccttattagt ggtggtggtg accgtgaaca cgctggtgct ggcacatggt 60
gttaccgaag tccactga 79

<210> 21

<211> 82

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic oligonucleotides used to join DNA fragments

<400> 21

gttcacggtc accaccacca ctcttcaaaa tgcggtctgt gcaccggttc tcagaaagaa 60
ggtctgcact acacctgctc tt 82

<210> 22
<211> 82
<212> DNA
<213> Artificial sequence

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<400> 22
ggtggatttt cagggctctgg aagtttttcc agaactggta ctgagagtac ggaagtga 60
aagagcaggt gtagtcaga cc 82

<210> 23
<211> 82
<212> DNA
<213> Artificial sequence

<220>
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<400> 23
ttccagaccc tgaaaatcca ccagcgtgtt caccatcacc attottacaa atgcggtctg 60
tgccaggaat tcttcggtct ga 82

<210> 24
<211> 93
<212> DNA
<213> Artificial sequence

<220>
<223> Synthetic oligonucleotides used to join DNA fragments

<400> 24
ggatccggat ccttattaag cagcgtgaac acgctgggtga ccgtccagac ggtagaaga 60
agagcagttg ttcagaccga agaattcctg gca 93

<210> 25
<211> 33
<212> DNA
<213> Artificial sequence

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<400> 25
catatgcata tggactacca ggtttcttct ccg 33

<210> 26
<211> 23
<212> DNA
<213> Artificial sequence

<220>
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<400> 26
gagtgggtgt ggtgaccgtg aac 23

<210> 27
<211> 42
<212> DNA
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<220>
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<400> 27
ggatccggat ccataaatat ggactaccag gtttcttctc cg 42

<210> 28
<211> 99
<212> DNA
<213> Artificial sequence

<220>
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<400> 28
ggatccggat ccataaatat gaaggctctcc gtggctgcc tctcctgcct catgcttggt 60
actgcccttg gatcgatgga ctaccagggt tcttctccg 99

<210> 29
<211> 48
<212> DNA
<213> Artificial sequence

<220>
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<400> 29
tctagatcta gattattaag cagcgtgaac acgctgggtga ccgtccag 48

<210> 30
<211> 41
<212> DNA
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<220>
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<400> 30
ggtaccggtta ccttattaag cagcgtgaac acgctgggtga c 41

<210> 31
<211> 157
<212> PRT
<213> Homo sapiens

<400> 31
Met Lys Val Ser Val Ala Ala Leu Ser Cys Leu Met Leu Val Thr Ala
1 5 10 15

Leu Gly Ser Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn
20 25 30

Tyr Tyr Thr Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala
35 40 45

Ala Tyr Lys Cys Gly Leu Cys Ala Ala Ala Gln Trp Asp Phe Gly Asn
50 55 60

Thr Met Cys Gln His Gln Arg Val His Gly His His His His Ser Tyr
65 70 75 80

Lys Cys Gly Leu Cys Thr Arg Ser Gln Lys Glu Gly Leu His Tyr Thr
85 90 95

Cys Ser Ser His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe
100 105 110

Gln Thr Leu Lys Ile His Gln Arg Val His Gly Gly Gly Gly Ser Tyr
115 120 125

Lys Cys Gly Leu Cys Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser
130 135 140

Ser Asn Arg Leu Asp Gly His Gln Arg Val His Ala Ala
145 150 155

<210> 32

<211> 19

<212> PRT

<213> Homo sapiens

<400> 32

Met Lys Val Ser Val Ala Ala Leu Ser Cys Leu Met Leu Val Thr Ala
1 5 10 15

Leu Gly Ser